

Communication

Stage-specific Expression of Ankyrin and SOCS Box Protein-4 (Asb-4) during Spermatogenesis

Soo-Kyoung Kim^{1,†}, Si Youn Rhim^{2,†}, Man Ryul Lee¹, Jong Soo Kim¹, Hyung Jun Kim^{1,3}, Dong Ryul Lee³, and Kye-Seong Kim^{1,*}

¹ Department of Anatomy and Cell Biology and Department of Biomedical Sciences, College of Medicine, Hanyang University, Seoul 133-791, Korea;

² Department of Surgery, Hanyang University, Seoul 133-791, Korea;

³ Fertility Center of CHA General Hospital, CHA Research Institute, Pochon CHA University, Seoul 135-081, Korea.

(Received November 5, 2007; Accepted December 13, 2007)

Members of the large family of Asb proteins are ubiquitously expressed in mammalian tissues; however, the roles of individual Asb and their function in the developmental testes have not been reported. In this report, we isolated a murine *Asb4* from mouse testis. Northern blot analysis revealed that *mAsb-4* was expressed only in testes and produced in a stage-specific manner during spermatogenesis. It was expressed in murine testes beginning in the fourth week after birth and extending into adulthood. Pachytene spermatocytes had the highest level of expression. Interestingly, the human homologue of *mAsb-4*, *ASB-4* (*hASB-4*) was also expressed in human testis. These results suggest that *ASB-4* plays pivotal roles in mammalian testis development and spermatogenesis.

Keywords: Ankyrin-Repeat; Asb-4; SOCS Box; Spermatogenesis.

Introduction

The suppressor of cytokine signaling (SOCS) family of proteins contributes to the negative regulation of cytokine signaling such as seen in the JAK-STAT pathway (Hilton et al., 1998; Kisseleva et al., 2002). Previous reports suggest that the SOCS box plays a role in ubiquitin-mediated proteasomal degradation via interaction with the elongin

B and elongin C complex (Chung et al., 2005; Kamura et al., 1998; Krebs and Hilton, 2000).

Ankyrin repeats occur in molecules with a wide variety of functions including receptors, cell-cycle regulators, membrane skeletal proteins, secreted proteins, tumor suppressors, and transcription factors (Breedon and Nasmyth, 1987; Kile et al., 2000; Zhang et al., 1999).

Spermatogenesis is a cyclic developmental process by which spermatogonial stem cells generate mature spermatozoa via a unique genetic and molecular program (Kho and Inaba, 2004; Rhee and Wolgemuth, 2002). Many germ cell-specific genes are involved in this developmental process, e.g., *mAsb-17*, as discussed earlier (Kim et al., 2004). The mechanism that regulates the genes involved is elaborate, and understanding of this finely tuned mechanism remains elusive.

Following on from our study of *mAsb-17*, we report that a member of the ankyrin repeat-containing SOCS box protein family, *Asb-4*, shows unique tissue- and stage-specific expression.

Materials and Methods

Mice gonads and testes Male ICR mice (5 d, 2, 4, 6, and 10 weeks old) and pregnant female ICR mice (12.5 dpc and 15.5 dpc) were purchased from Daehan Biolink Co. Ltd. (Korea). The procedures for animal care and management was followed by SOP of Hanyang University, College of Medicine, Animal Care and Use Committee.

Collecting gonadal germ cells To collect germ cells, gonads were dissected and removed from 12.5 dpc and 15.5 dpc embryos with fine hypodermic needles and forceps. They were washed briefly in calcium- and magnesium-free phosphate-buffered solu-

[†] These authors contributed equally to this work.

* To whom correspondence should be addressed.
Tel: 82-2-2220-0607; Fax: 82-2-2281-7841
E-mail: ks66kim@hanyang.ac.kr

tion and incubated in an EDTA solution (20 mg EDTA, 800 mg NaCl, 20 mg KCl, 115 mg Na₂HPO₄, 20 mg KH₂PO₄ in 100-ml distilled water) for 20 min at room temperature. They were then washed in Dulbecco's modified Eagle's medium (DMEM) [10% fetal bovine serum (FBS)] and gently disrupted using a fine needle. The released germ cells were collected with a fine-drawn Pasteur pipette.

Isolation of spermatogenic cell populations Mixed populations of spermatogenic cells were obtained from testes of 10-weeks-old male mice using the collagenase dissociation method (Bellve et al., 1977). Purified populations of spermatogenic cells were isolated following collagenase treatment of testes and trypsin digestion of isolated seminiferous tubules using unit gravity sedimentation in a bovine serum albumin gradient (Bellve et al., 1977; Romrell et al., 1976). Tubes containing each cell population (pachytene spermatocyte (PS), > 90% pure; round spermatid (RS), > 90% pure; and condensing spermatid (CS)-residual body mixture) were selected by phase contrast microscopy.

Isolation and analysis of human testicular tissue from male infertility patients Testicular tissues were obtained from male infertility patients using the multiple testicular sperm extraction (TESE)-ICSI program. When sperm or round spermatids were absent from dissected samples, the remaining testicular tissues were donated for experiment with informed consent (Lee et al., 2006). This study was approved by the Institutional Review Board of the CHA General Hospital (Korea). To identify the existence of germ cells in the human testis samples from the male infertility patients, we investigated the expression of the specific marker genes for transition protein (TP)-1 and protamine. After confirming the presence of germ cells, RT-PCR analysis of *hASB-4* was performed.

STO and mouse ESC Culture Mouse STO fibroblast feeder cells [CRL-1503; American Type Culture Collection (ATCC), USA] were grown in DMEM (GIBCO/BRL, USA) supplemented with 10% FBS (HyClone, USA) and 1% penicillin-streptomycin (GIBCO/BRL, USA). These cells were mitotically inactivated with 10 µg/ml mitomycin-C (Sigma, USA) for 1.5 h. HS-3 mouse ESC were grown under standard conditions. Both cell types were grown in 5% CO₂, 95% air and were routinely passaged every 4–5 days.

RT-PCR for expression analysis Total cell RNA was isolated from mouse gonads, testes, STO, and mESCs using TRIzol (Invitrogen, USA). Whole mouse testes were obtained from mice at 5 days, and 2, 4, 6, and 10 weeks. PS, RS, and CS were separated from whole testes of 10-weeks-old mice by STA-PUT. One microgram of total RNA from each population was treated with DNase I (Sigma, USA). cDNA synthesis from total RNA was performed according to the protocol of the Super Script Preamplification system (Invitrogen, USA). After incubation of 1 µg total RNA with 0.5 µg oligo (dT)₁₂₋₁₈ primer at 70°C for 10

min, the reaction was carried out in the 5X first strand buffer (containing 250 mM Tris-HCl, 375 mM KCl, and 15 mM MgCl₂) from the kit, 10 mM DTT (dithiothreitol) and 0.5 mM dNTPs in a final volume of 20 µl at 42°C for 2 min. Then SuperScript™ II (Invitrogen, USA) was added and incubated for 50 min at 42°C. After the reaction, the enzyme was denatured at 70°C for 15 min. Primers used for amplification of *mAsb-4* were 5'-CAAAGCT-CAACTGCTACTCC-3' (forward) and 5'-CAG-CTGGTAGCA-GATCTCA-3' (reverse; product 539 bp) (GenBank accession number: *mAsb-4*; BC046819). Amplification was achieved by 28 cycles of 94°C for 35 s, 55°C for 40 s, and 72°C for 45 s followed by a final incubation for 7 min at 72°C. As a loading control, the same amounts of cDNA templates were used to amplify *G3PDH*. The oligonucleotide primers used for amplification of *mG3PDH* were 5'-ACTGGTGCTGCC-AAGGCT-GT-3' (forward) and 5'-CGGCATCGAAGGTGG-AAGAG-3' (reverse; product 262 bp).

To identify the expression of specific marker genes in the testes from male infertility patients, we investigated the expression of the TP-1 and protamine genes. Primers used for amplification were 5'-gtcaagagag gtggcagcaa -3' (forward) and 5'-tcacaagtgaggcggtaat -3' (reverse; product 104 bp) (GenBank accession number: NM_003284) and 5'-cgaggtgtac aggcagcagt-3' (forward) and 5'-gctctctcg agagcagtg-3' (reverse; product 145 bp) (GenBank accession number: NM_002761), respectively. Amplification was achieved by 30 cycles of 94°C for 35 s, 55°C for 30 s, and 72°C for 30 s followed by a final incubation for 5 min at 72°C.

Northern blot analysis Expression of *mAsb-4* mRNA from mouse tissue was analyzed by Northern blot analysis. Mouse multiple tissue blot was purchased from Seegene (Seegene, Korea). For the stage-specific mouse blot, RNA samples (20 µg each) of STO, 12.5 dpc, 15.5 dpc, 5 days, 2 weeks, 4 weeks, 6 weeks, 10 weeks, pachytene spermatocyte, round spermatid, condensing spermatid and mESCs were electrophoresed and blotted onto Hybond™-N+ membranes (Amersham Biosciences, UK). The immobilized nucleic acids were hybridized with the *mAsb-4* probe labeled by random priming with [α -³²P] dCTP (Amersham Biosciences, UK). Hybridization was performed in a bag containing ExpressHyb™ Hybridization solution (Clontech, USA) at 68°C overnight. The hybridized membrane was washed at room temperature in 2× SSC and 0.1% SDS, then in 0.1× SSC and 0.1% SDS, and exposed to Kodak X-ray film with an intensifying screen for 5 h at -70°C.

Results

Alignment of amino acid sequences for mouse, and human *Asb-4* Mouse *Asb-4* has an open reading frame of 1278 bp encoding a putative protein of 426 amino acids. Figure 1A shows the amino acid sequences of *Asb-4* family members searched in GenBank aligned using the MegAlign with DNASTAR program. The amino acid sequence of

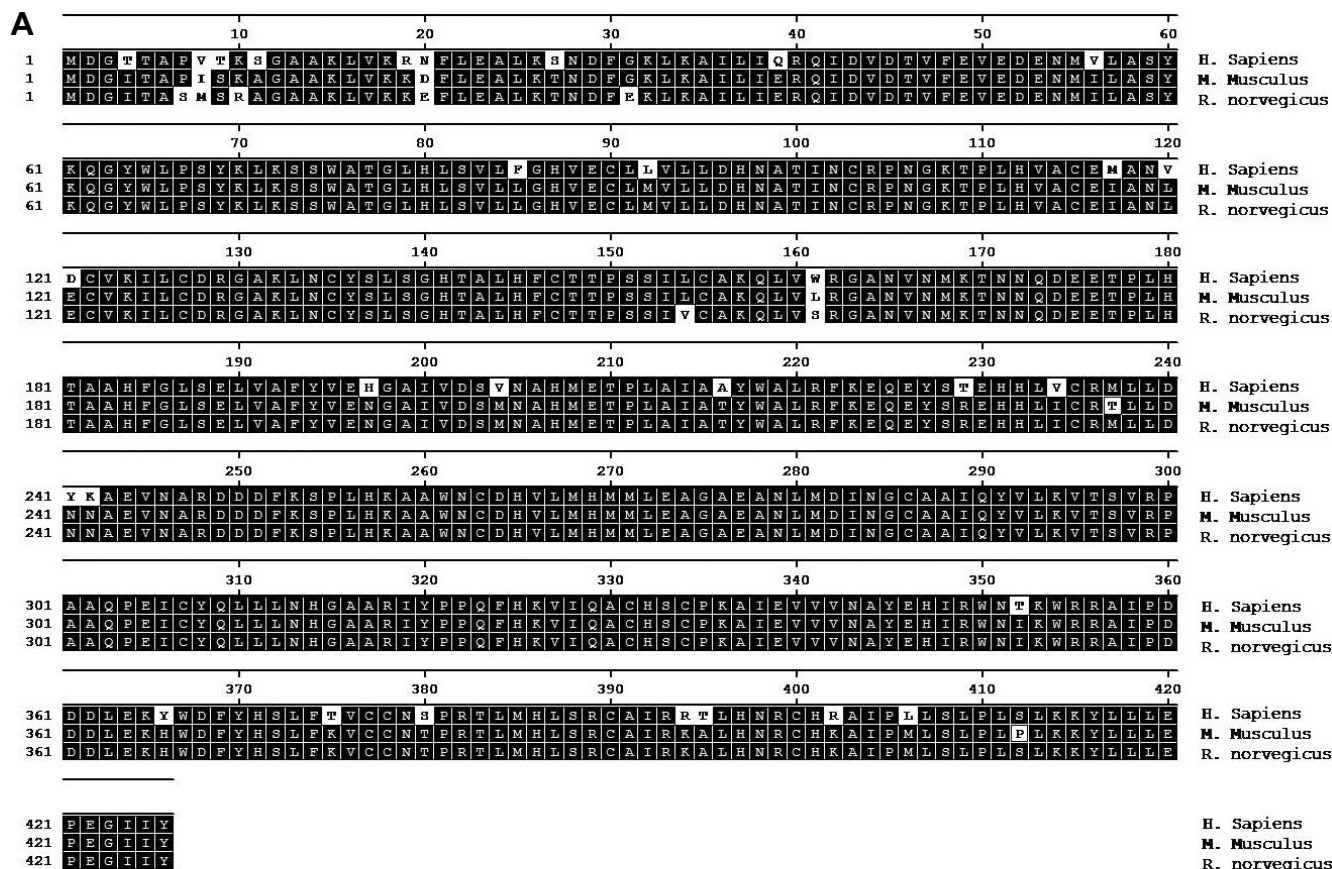


Fig. 1. **A.** Alignment of amino acid sequences of mouse *Asb-4* (NP_075535), human *ASB-4* (NP_057200), Rat *Asb-4* (NP_001019489) using MegAlign with DNASTAR program. **B.** Domain of mouse *Asb-4* contains six ankyrin repeats (ANK: amino acids 74-103, 106-135, 139-168, 174-203, 207-247, and 251-280) and one SOCS box (371-423).

mAsb-4 has 92% and 97% identity to that of humans and rat respectively (Table 1). SMART (<http://smart.embl-heidelberg.de>) showed that the deduced amino acid sequence of the *mAsb-4* contains six ankyrin repeat domains (74–103, 106–135, 139–168, 174–203, 207–247 and 251–280 amino acids, respectively), and one SOCS box domain between amino acids 371 and 423 (Fig. 1B).

Expression of *Asb-4* in mouse and human tissue To characterize the expression of *mAsb-4* in various tissues, we performed a Northern blot analysis (Fig. 2A). Total RNA was prepared from mouse brain, heart, lung, liver, spleen, kidney, stomach, small intestine, skeletal muscle, thymus, testis, non-pregnant uterus, and placenta. Strikingly, *mAsb-4* expression was seen only in the testis. We performed RT-PCR using a pair of primers designed from the sequence of a cDNA library clone (GenBank acces-

Table 1. The nucleotide and amino sequences of *mAsb-4* and other members of the *Asb-4* family.

Mouse (<i>mAsb-4</i>)	Nucleotide sequence identity (%)	Amino acid sequence identity (%)
Human	85	92
Rat	93	97

sion number: BC046819; Fig. 2B) to further investigate expression in various spermatogenic cells. PCR showed that *mAsb-4* was expressed after the fourth week of mouse testis development (Fig. 2B; lanes 4–6). In addition, Northern blot analysis revealed that the *mAsb-4* transcript of about 3 kb length was most strongly expressed in pachytene spermatocytes and was down regulated during further progression into condensing spermatids (Fig. 2C).

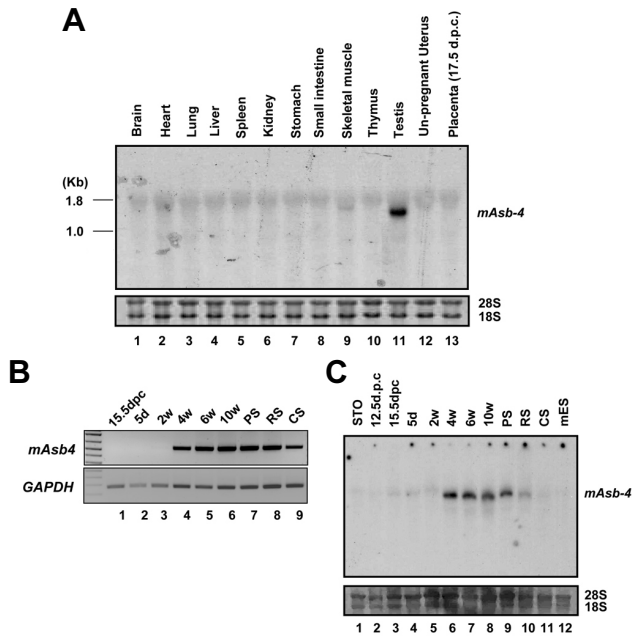


Fig. 2. **A.** Expression of *mAsb-4* mRNA in mouse organs. Total RNA was prepared from mouse brain (lane 1), heart (lane 2), lung (lane 3), liver (lane 4), spleen (lane 5), kidney (line 6), stomach (lane 7), small intestine (lane 8), skeletal muscle (lane 9), thymus (lane 10), testis (lane 11), non-pregnant uterus (lane 12), and placenta (lane 13). Expression was seen only in testis. 28s/18s RNAs were used as loading controls. **B.** *mAsb-4* expression during mouse testis development and spermatogenesis, shown by RT-PCR. Lane 1 is a testicular sample from 15.5 dpc fetal gonads and lanes 2–6 are testicular samples from 5-day-old, 2-, 4-, 6-, and 10-weeks-old mice. In addition, cDNA samples from pachytene spermatocytes (PS), round spermatids (RS), and condensing spermatids (CS) were used. To assess RNA integrity, a *GAPDH* control was included. **C.** Northern blot hybridization of *mAsb-4* mRNA from 12.5 and 15.5 dpc fetal gonads (lanes 2, 3), 5-day-old, 2-, 4-, 6-, and 10-weeks-old mouse testis, pachytene spermatocytes (PS), round spermatids (RS), and condensing spermatids (CS) during spermatogenesis. mRNA from mouse embryonic stem cells (mESC) and STO cells was used as control. 28s/18s RNAs were served as loading controls.

As there were no earlier reports of expression of *hASB-4* in human testicular samples we obtained human testicular samples from male infertility patients and tested for the presence of germ cells using the germ cell marker genes for TP-1 and protamine, and, examined the expression of *hASB-4* in the same samples. Interestingly, *hASB-4* was expressed only in the samples containing germ cells (Fig. 3; lane 2).

Discussion

Cytokines regulate fundamental biological processes such

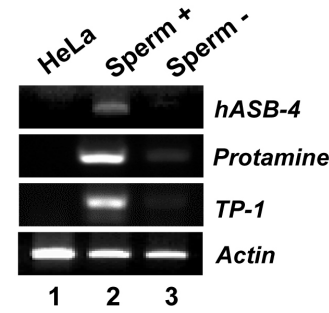


Fig. 3. RT-PCR analysis of *hASB-4* and germ-cell-specific marker genes in testicular tissues isolated from male infertility patients. Expression of TP-1 and protamine genes indicates the presence of germ cells. *hASB-4* was uniquely expressed in germ cell-containing samples. HeLa cells served as negative control. To determine RNA integrity, an actin positive control was included.

as immunity, wound healing, and hematopoiesis by interacting with receptors on cell surfaces (Ochoa et al., 2000). Negative feedback is another important mechanism regulating signal transduction by a wide variety of extracellular signals. Suppressors of cytokine signaling (SOCS) proteins are known as negative regulators of signal transduction. It is assumed that the SOCS box plays a role as a negative regulator in signal transduction, perhaps by inhibiting kinase activity. Another possibility is that the SOCS box acts as an adaptor module or regulates aspects of protein behavior such as intracellular location or stability (Hilton et al., 1998).

Very little is known about the ASB subfamily. Testes lacking *Asb-1* show a reduction of spermatogenesis with less complete filling of the seminiferous tubules (Kile et al., 2000; 2001). Exogenous expression of *Asb-2* in myeloid leukemia cells resulted in growth inhibition and chromatin condensation (Guibal et al., 2002). *Asb-6* interacts with APS to enable recruitment of elongins B and C to the insulin receptor-signaling complex in adipocytes. In sum, it is thought that ASB family members have pleiotropic functions through protein-protein interactions.

In addition to expression of *mAsb-17* as we described earlier, *mAsb-4* expression has been reported to be restricted to mouse testis (lanes 8, 11); we confirmed this result here (Fig. 2A). In addition, we found that *mAsb-4* expression differed noticeably during spermatogenesis. Mice attain puberty at 5–6 weeks. The testes descend into the scrotum and produce sperm after the fifth week and male mice develop sexual maturity between 6 and 8 weeks of age. The pachytene phase of meiosis has a fixed duration of over a week for each particular species in pachytene spermatocytes (meiotic cells). Genetic recombination also occurs during this period. After the second meiosis, round spermatids (early haploid cells) undergo a series of shape changes, and condensing spermatids (late

haploid cells) also undergo marked transformations ultimately producing species-specifically shaped spermatozoa, in a process referred to as spermiogenesis. Global inhibition of the initiation of mRNA translation occurs in pachytene spermatocytes and round spermatids (Park et al., 2001). *mAsb-4* begins to be expressed near puberty and is continuously expressed thereafter. In spermatogenic cells, pachytene spermatocytes show the strongest expression of *mAsb-4*, while in the next stage of spermatogenic cells, round spermatids have weaker expression levels (Fig. 2C). Human testicular samples containing developing germ cells also showed expression of human *ASB-4*. Our findings imply that this unique gene is closely associated with testes development and spermatogenesis.

Acknowledgment This work was supported by the research fund of Hanyang University (HY-2004) to KSK.

References

- Bellve, A.R., Millette, C.F., Bhatnagar, Y.M., and O'Brien, D.A. (1977). Dissociation of the mouse testis and characterization of isolated spermatogenic cells. *J. Histochem. Cytochem.* *25*, 480–494.
- Breeden, L., and Nasmyth, K. (1987). Similarity between cell-cycle genes of budding yeast and fission yeast and the Notch gene of *Drosophila*. *Nature* *329*, 651–654.
- Chung, A.S., Guan, Y.J., Yuan, Z.L., Albina, J.E., and Chin, Y.E. (2005). Ankyrin repeat and SOCS box 3 (ASB3) mediates ubiquitination and degradation of tumor necrosis factor receptor II. *Mol. Cell. Biol.* *25*, 4716–4726.
- Guibal, F.C., Moog-Lutz, C., Smolewski, P., Di, Gioia, Y., Darzynkiewicz, Z., Lutz, P.G., and Cayre, Y.E. (2002). ASB-2 inhibits growth and promotes commitment in myeloid leukemia cells. *J. Biol. Chem.* *277*, 218–224.
- Hilton, D.J., Richardson, R.T., Alexander, W.S., Viney, E.M., Willson, T.A., Sprigg, N.S., Starr, R., Nicholson, S.E., Metcalf, D., and Nicola, N.A. (1998). Twenty proteins containing a C-terminal SOCS box form five structural classes. *Proc. Natl. Acad. Sci. USA* *95*, 114–119.
- Kamura, T., Sato, S., Haque, D., Liu, L., Kaelin, W.G., Jr., Conaway, R.C., and Conaway, J.W. (1998). The Elongin BC complex interacts with the conserved SOCS-box motif present in members of the SOCS, ras, WD-40 repeat, and ankyrin repeat families. *Genes Dev.* *12*, 3872–3881.
- Kho, K.H., and Inaba, K. (2004). Expression of 17beta-hydroxysteroid dehydrogenase in testis of the ascidian *Ciona intestinalis* [corrected]. *Mol. Cells* *18*, 171–176.
- Kile, B.T., Viney, E.M., Willson, T.A., Brodnicki, T.C., Cancilla, M.R., Herlihy, A.S., Croker, B.A., Baca, M., Nicola, N.A., Hilton, D.J., et al. (2000). Cloning and characterization of the genes encoding the ankyrin repeat and SOCS box-containing proteins Asb-1, Asb-2, Asb-3 and Asb-4. *Gene* *258*, 31–41.
- Kile, B.T., Metcalf, D., Mifsud, S., DiRago, L., Nicola, N.A., Hilton, D.J., and Alexander, W.S. (2001). Functional analysis of Asb-1 using genetic modification in mice. *Mol. Cell. Biol.* *21*, 6189–6197.
- Kim, K.S., Kim, M.S., Kim, S.K., and Baek, K.H. (2004). Murine Asb-17 expression during mouse testis development and spermatogenesis. *Zygote* *12*, 151–156.
- Kisseleva, T., Bhattacharya, S., Braunstein, J., and Schindler, C.W. (2002). Signaling through the JAK/STAT pathway, recent advances and future challenges. *Gene* *285*, 1–24.
- Krebs, D.L., and Hilton, D.J. (2000). SOCS: physiological suppressors of cytokine signaling. *J. Cell Sci.* *113 (Pt 16)*, 2813–2819.
- Lee, D.R., Kim, K.S., Yang, Y.H., Oh, H.S., Lee, S.H., Chung, T.G., Cho, J.H., Kim, H.J., Yoon, T.K., and Cha, K.Y. (2006). Isolation of male germ stem cell-like cells from testicular tissue of non-obstructive azoospermic patients and differentiation into haploid male germ cells *in vitro*. *Hum. Reprod.* *21*, 471–476.
- Ochoa, G.C., Slepnev, V.I., Neff, L., Ringstad, N., Takei, K., Daniell, L., Kim, W., Cao, H., McNiven, M., Baron, R., et al. (2000). A functional link between dynamin and the actin cytoskeleton at podosomes. *J. Cell Biol.* *150*, 377–389.
- Park, C., Choi, W.S., Kwon, H., and Kwon, Y.K. (2001). Temporal and spatial expression of neurotrophins and their receptors during male germ cell development. *Mol. Cells* *12*, 360–367.
- Rhee, K., and Wolgemuth, D.J. (2002). Tcp10 promoter-directed expression of the Nek2 gene in mouse meiotic spermatocytes. *Mol. Cells* *13*, 85–90.
- Romrell, L.J., Bellve, A.R., and Fawcett, D.W. (1976). Separation of mouse spermatogenic cells by sedimentation velocity. a morphological characterization. *Dev. Biol.* *49*, 119–131.
- Zhang, J.G., Farley, A., Nicholson, S.E., Willson, T.A., Zugaro, L.M., Simpson, R.J., Moritz, R.L., Cary, D., Richardson, R., Hausmann, G., et al. (1999). The conserved SOCS box motif in suppressors of cytokine signaling binds to elongins B and C and may couple bound proteins to proteasomal degradation. *Proc. Natl. Acad. Sci. USA* *96*, 2071–2076.